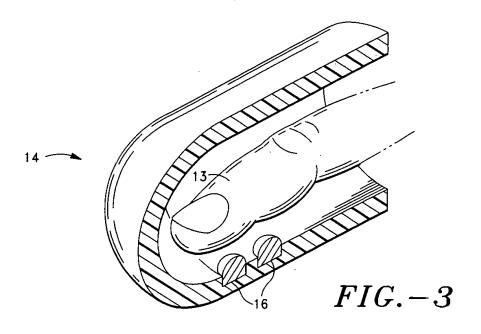
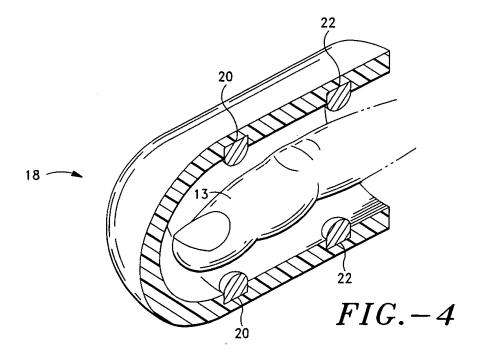
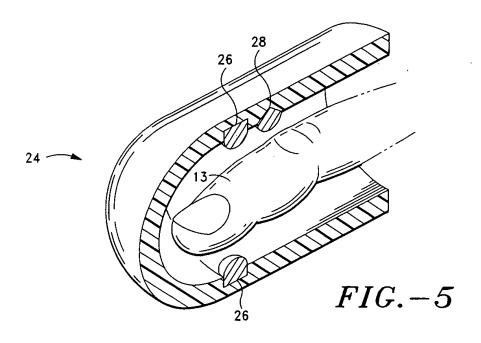
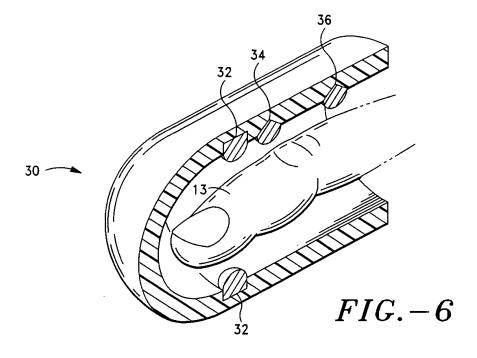


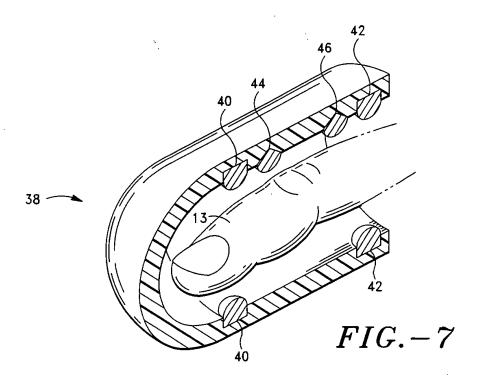
FIG.-2

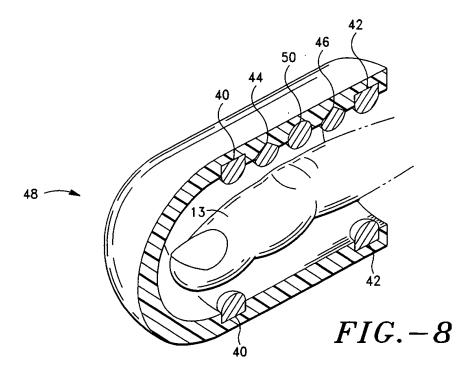












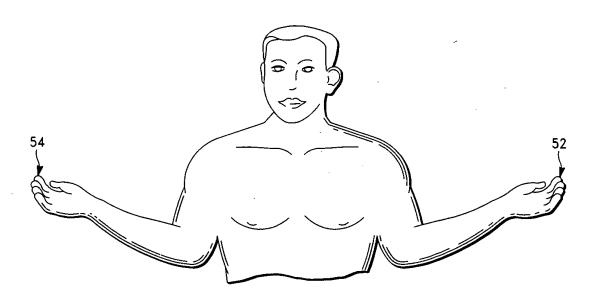


FIG.-9

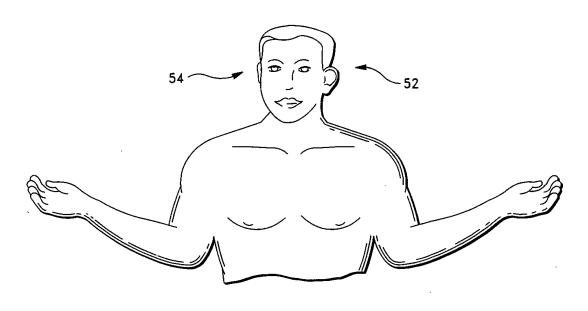


FIG. - 10

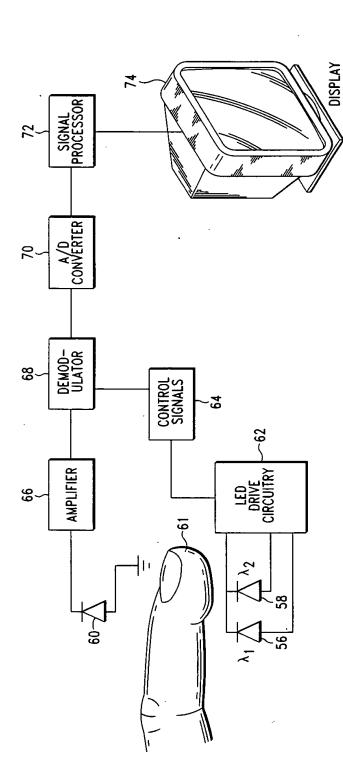


FIG.-11

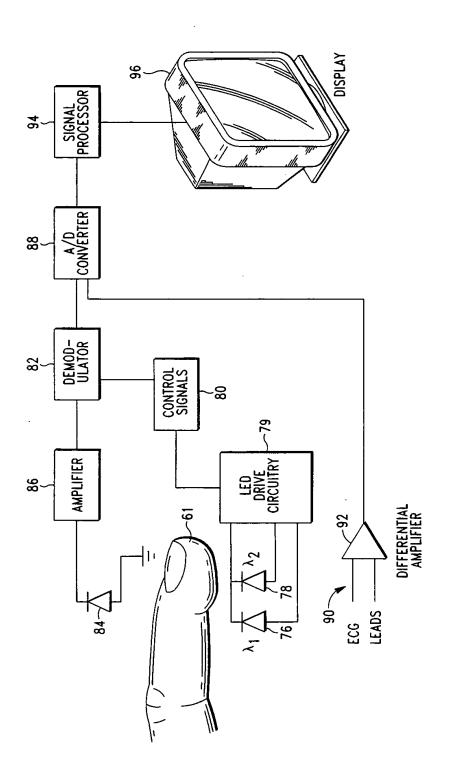
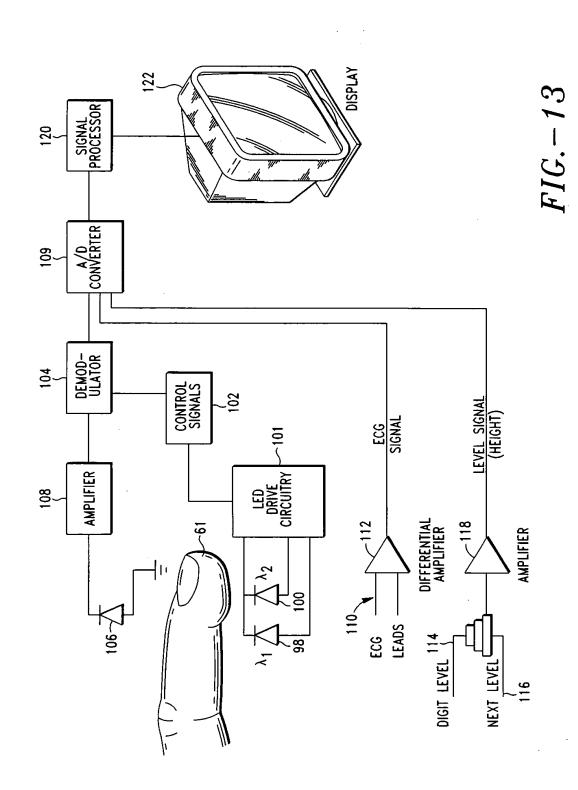


FIG.-12



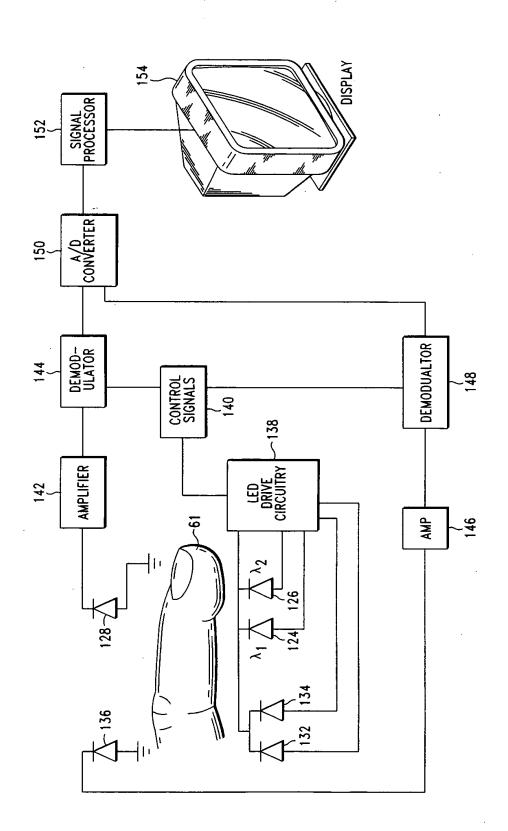
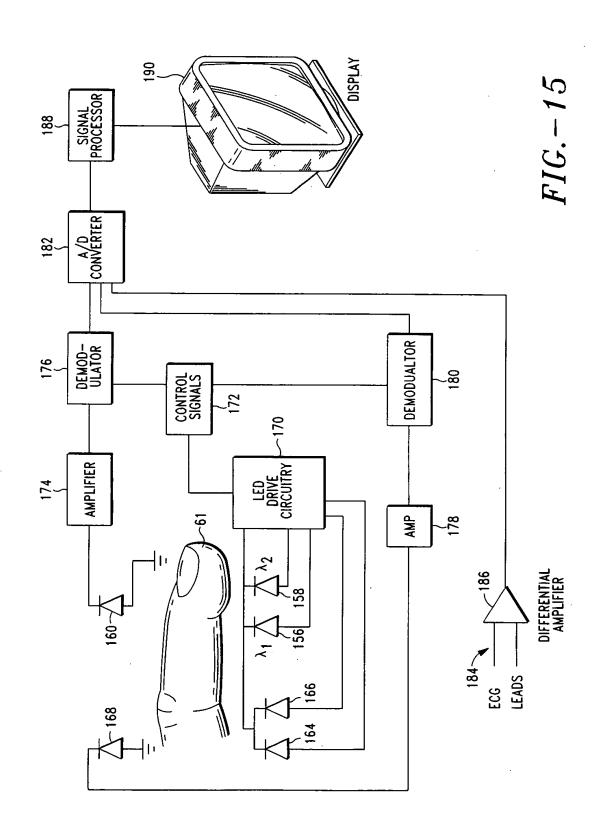


FIG.-14



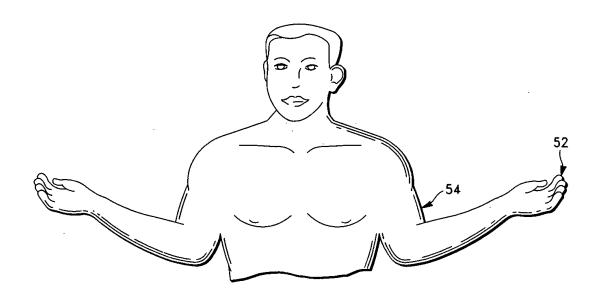


FIG.-16

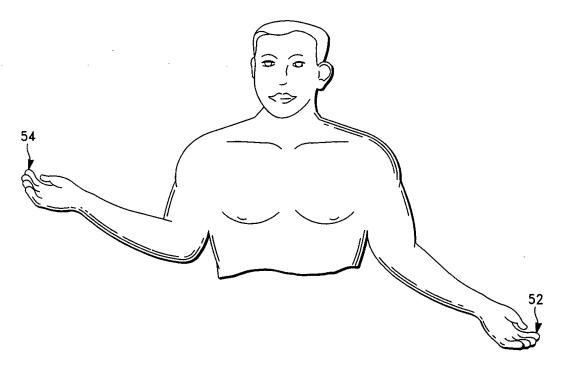
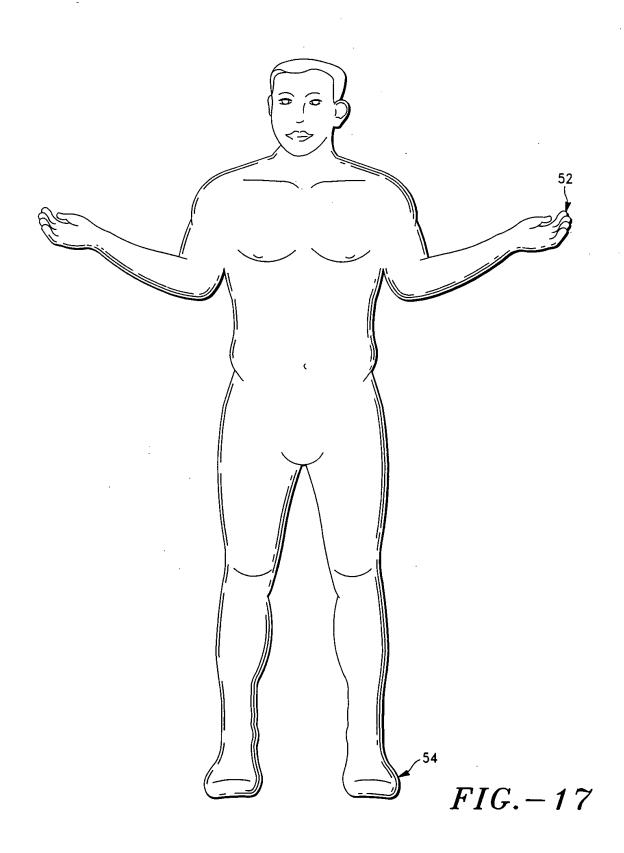
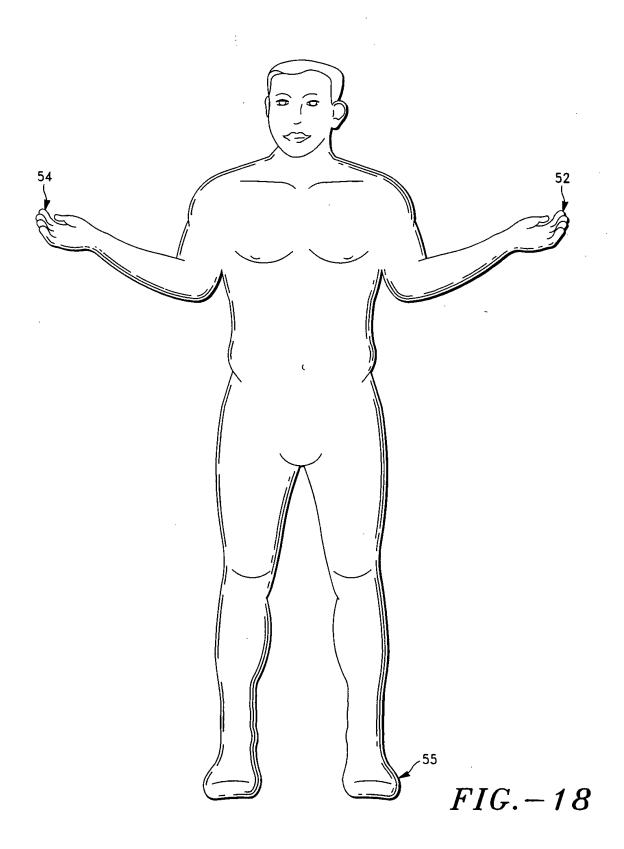


FIG.-19





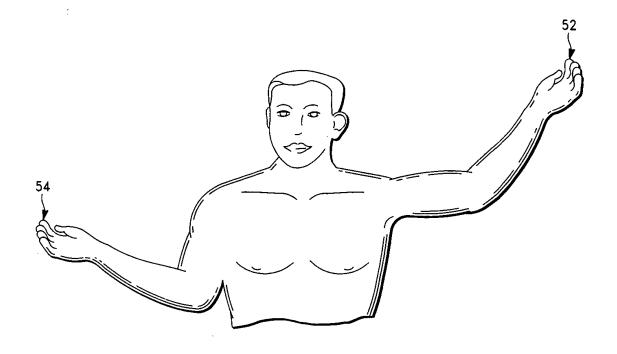
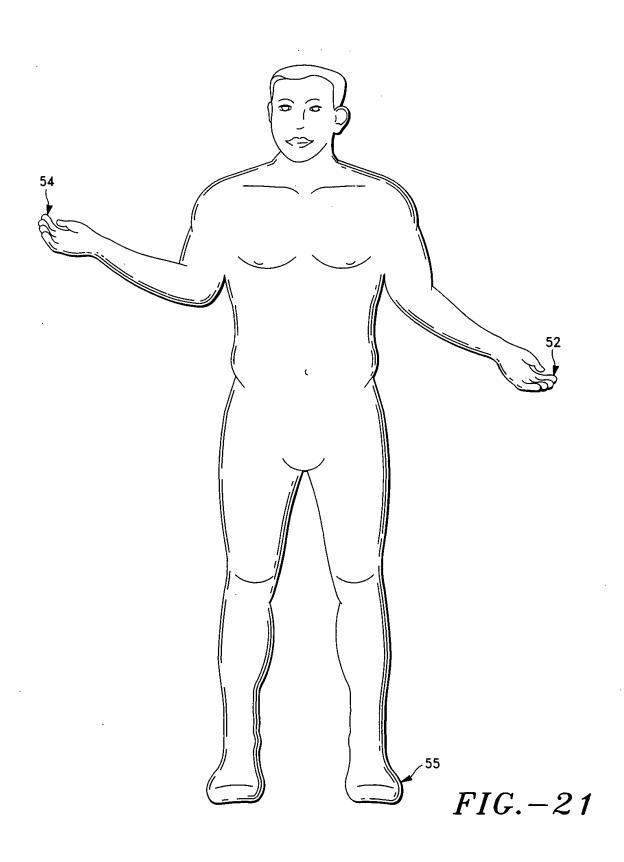


FIG.-20



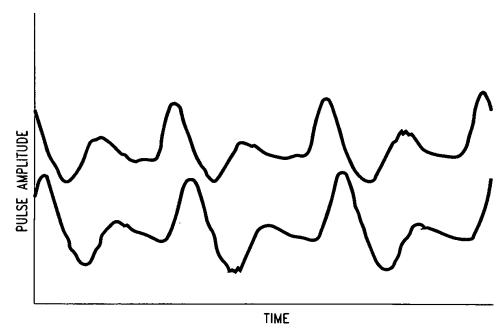


FIG.-22

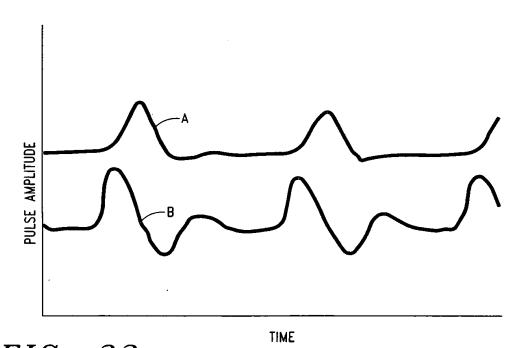


FIG.-23

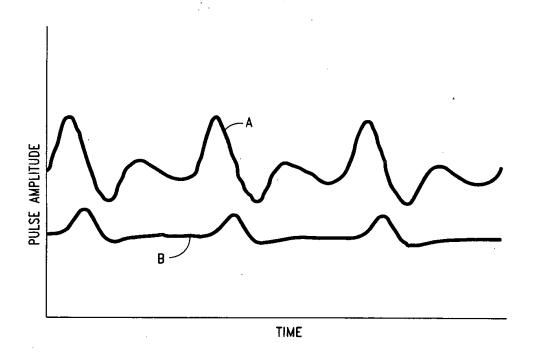


FIG.-24

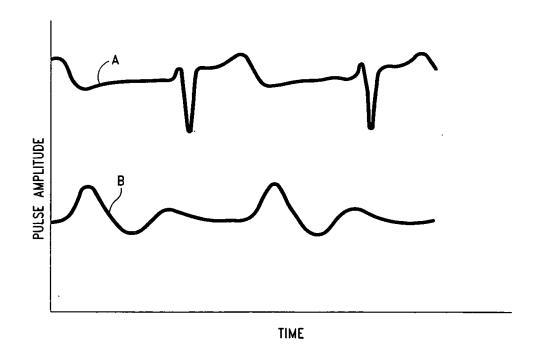


FIG.-25

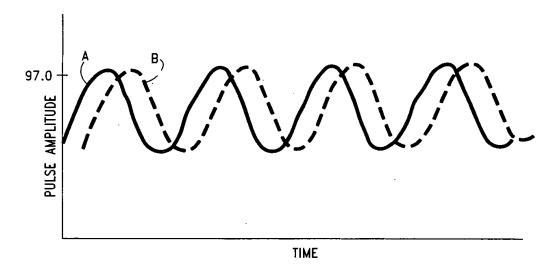


FIG.-26

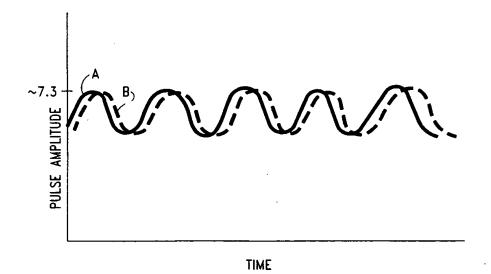
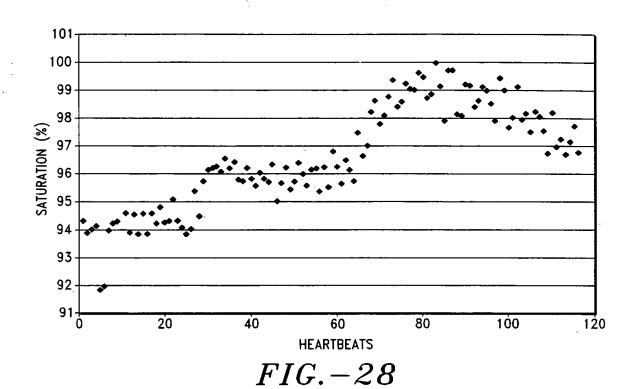
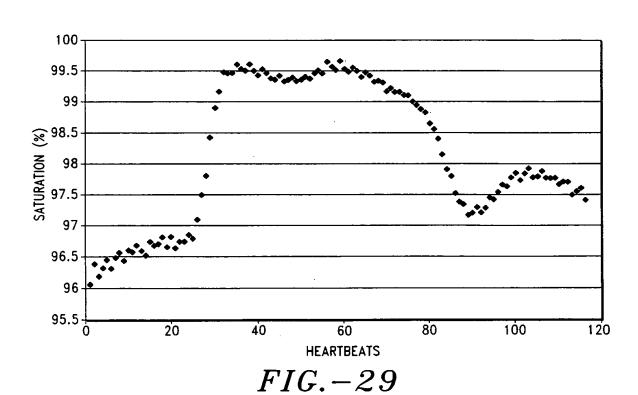
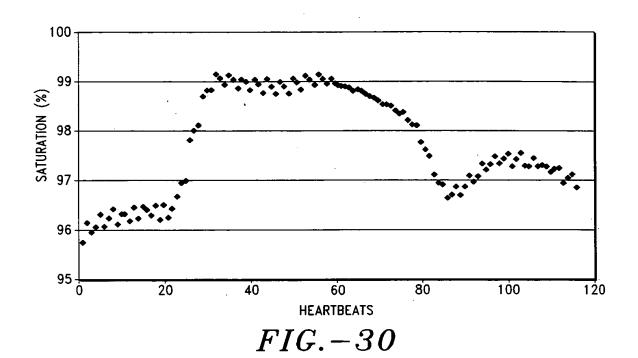
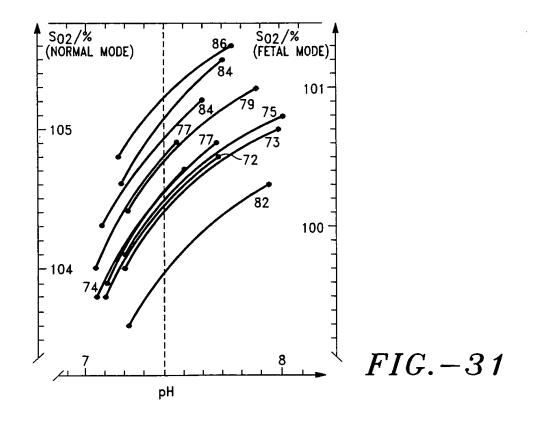


FIG.-27









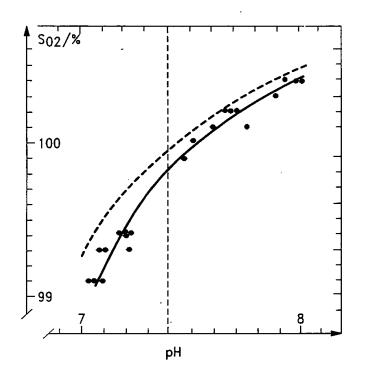


FIG.-32

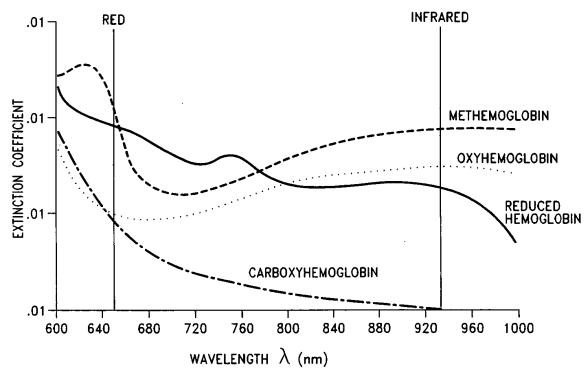


FIG.-34

```
Computational Algorithm for Determination of Hemoglobin Concentration
000000
           A is the measured absorbance
          A1 is the absorbance after dividing out extinction coefficients and correcting for saturation A2,A3, ... will be the absorbances at different path lengths, created by multiplying by constants A1,A3, ... and L2,L3, etc.
           constant M2=0.9 constant M3=0.8
           constant M4=0.7
           constant M5=0.6
           constant M6=0.5
           constant M7=0.4
           constant M8=0.3
C
CCC
           read in the value for hemoglobin absorbance and a value k
           representing the extinction coefficient for the wavelength and
           the oxygen saturation
           Begin
           Read, A
           Read, k
A1:=A/k
           A2:=A1*M2
           A3:=A1*M3
A4:=A1*M4
           A5:=A1*M5
           A6:=A1*M6
           A7:=A1*M7
           A8:=A1*M8
C
           k1234 = log(A1) * log(A2) - log(A3) * log(A4)

k5678 + log(A5) * log(A6) - log(A7) * log(A8)
           kd:=[\log(A1*A2) - \log(A3*A4)] / [\log(A5*A6) - \log(A7*A8)]
ç
           combine all the A terms that occur as coefficients,
           kAc := log(A2/A1) - log(A3/A1) - log(A4/A1) - [(kd * log(A5/A1)] -
                     - [kd * log(A6/A1) + [kd * log(A7/A1)] + [kd * log(A8/A1)]
C
           combine all the A terms that occur alone
           kAa := - [log(A_3/A_1) * log(A_4/A_1)] -
                              - kd * [log(A_5/A_1) * log(A_6/A_1)] + kd* [(log(A_7/A_1) * log(A_8/A_1)]
C
          k_{1234} - (k_d * k_{5678}) = kig(L) * k_{Ac} + k_{Aa} \log(L) = [k_{1234} - (k_d * k_{5678}) - k_{Aa}] / k_{Ac} L = antilog \{ [k_{1234} - (k_d * k_{5678}) - k_{Aa}] / k_{Ac} \}
use EXP or antilog function
L = EXP([k_{1234} - (k_d * k_{5678}) - k_{Aa}] / k_{Ac})
L is the path length
C
С
C
C
           C is the concentration of hemoglobin C = A1 / L
C
           END
```

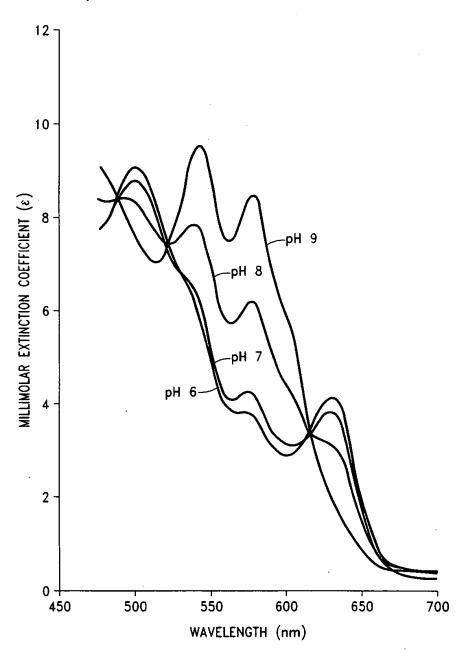


FIG.-35

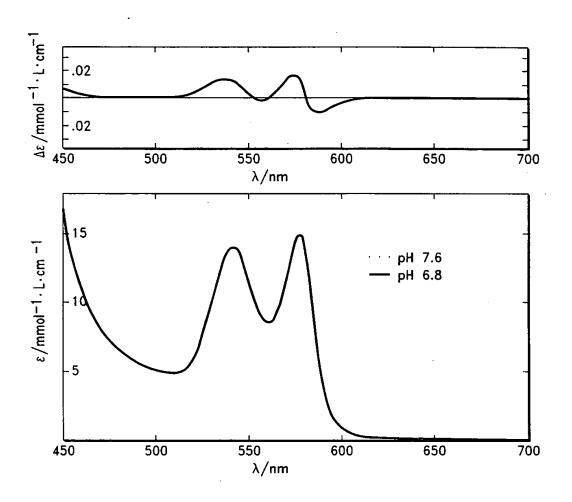


FIG.-36

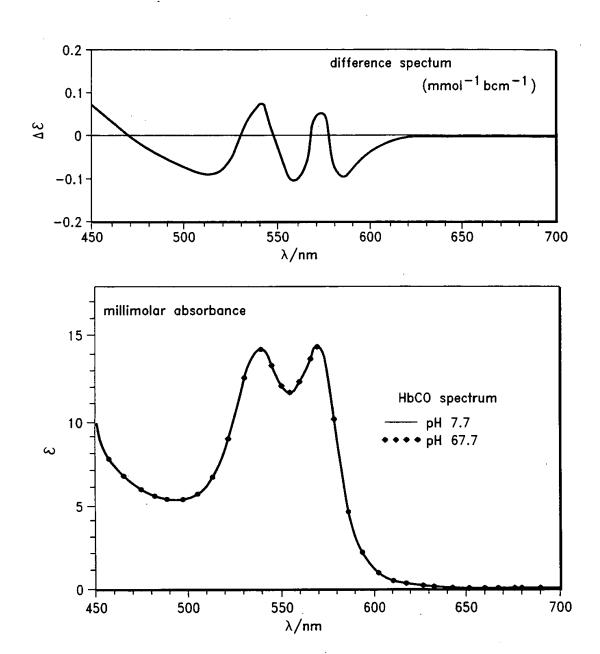
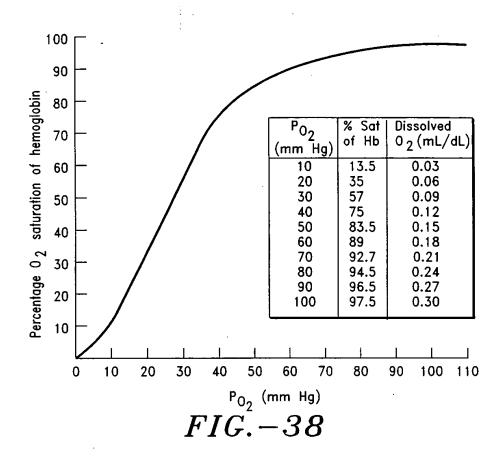


FIG.-37



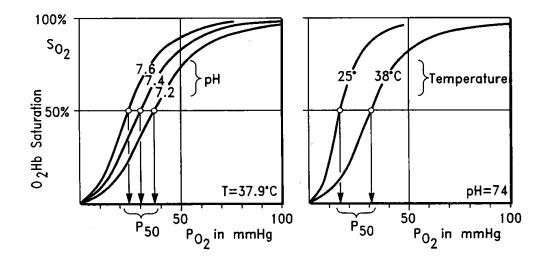


FIG.-39

